FISH GENETICS

Robb F. Leary, Ph.D

Division of Biological Sciences University of Montana Missoula, Montana 59812

U.S. Fish & Wildlife Service National Conservation Training Center Course #FIS1102

Table of Contents

Chapter I: Molecular Genetics and Cytogenetics	I-1
Introduction	I-3
What is DNA?	I-3
What is a gene?	I-6
What is a chromosome?	I-6
Protein Construction	I-9
What is RNA?	I-9
Genetic code	I-14
Summary	I-14
Cell Division	I-16
DNA replication	I-16
Mitosis	I-18
Meiosis	I-22
Contrast of mitosis and meiosis	I-27
Triploids and Gynogenetic Diploids	I-28
Problems and Discussion I	I-29
Chapter II: Introductory Genetics	II-1
Genetic variation	II-2
Commonly used terms	
Inheritance of single gene traits	
Mendelian ratios	
Inheritance of multiple gene traits	
Heritability	II-11
Problems and Discussion II	II-16
Chapter III: Introductory Population Genetics and Evolution	III-1
Evolution	
The nonevolving or equilibrium population	III-2
Hardy-Weinberg equilibrium model	
Mutation	III-7
Natural selection	III-8
Migration or gene flow	. III-11
Genetic drift	. III-13

Related topics	III-16
Founder effect	III-16
Bottleneck	III-17
Population genetic structure	
Problems and Discussion III	III-20
Chapter IV: Genetic Variation and Divergence	IV-1
Estimating Genetic Variation and Divergence	IV-2
Historical views	IV-2
Problems	IV-2
Solutions	IV-2
Protein electrophoresis	IV-3
Estimating allele frequencies	
Estimating genetic variation within populations	
Proportion of polymorhic loci	
Average expected heterozygosity	
Average number of alleles per locus	
Estimating genetic divergence among populations	
Similarity or distance estimates	
Principle components\discriminant function	IV-8
Partitioning genetic variation	
DNA analysis	
Restriction enzymes	
Polymerase chain reaction (PCR)	
Random amplified polymorphic DNA (RAPD)	
Microsatellites	
Problems and Discussion IV	IV-15
Data Interpretation	IV-18
Taxonomic questions	IV-18
Population genetic structure of a species	IV-22
Neutrality	
Importance of Genetic Variation	IV-26
Long term	
Short term	IV-26

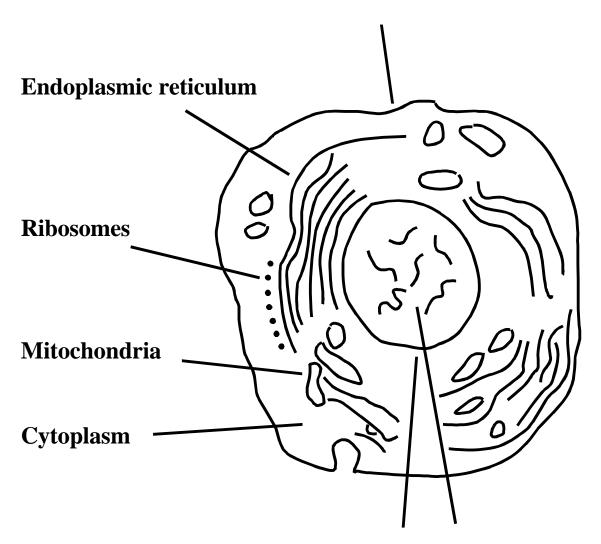
Causes of the loss of genetic variation IV-31 Inbreeding IV-31 Directional selection IV-33 Genetic drift IV-33 Genetic drift IV-33 Establishing a broodstock IV-33 Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Seasons for introductions IV-2 Reasons for introductions IV-2 Hybridization/Introgression IV-3 Indirect genetic changes due to reduced population size or changes in selection pressures IV-7 Stage II IV-7 Stage II IV-7 Stage III IV-7 Founding new populations of threatened fishes IV-7 LITERATURE CITED IVI	Maintaining Genetic Variation	IV-31
Directional selection IV-33 Genetic drift IV-33 Establishing a broodstock IV-33 Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-2 Reasons for introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	Causes of the loss of genetic variation	IV-31
Directional selection IV-33 Genetic drift IV-33 Establishing a broodstock IV-33 Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-2 Reasons for introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	Inbreeding	IV-31
Establishing a broodstock IV-33 Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	Genetic drift	IV-33
Number of founders IV-33 General IV-33 Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	Establishing a broodstock	IV-33
Considerations from spawning time distributions IV-33 Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	General	IV-33
Threatened and endangered species IV-37 Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Perpetuating a broodstock IV-38 Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Transferring a broodstock IV-42 Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Monitoring a broodstock IV-42 Chapter V: Fish Introductions V-1 Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI		
Genetic Impacts of Fish Introductions V-2 Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	Chapter V: Fish Introductions	V-1
Reasons for introductions V-2 Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	<u>-</u>	
Hybridization/Introgression V-3 Indirect genetic changes due to reduced population size or changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	<u>-</u>	
Indirect genetic changes due to reduced population size or changes in selection pressures		
changes in selection pressures V-6 Proper Use of Fish Introductions in Population Restoration V-7 Stage I V-7 Stage II V-7 Stage III V-7 Founding new populations of threatened fishes V-7 LITERATURE CITED VI	·	
Stage II		V-6
Stage II	Proper Use of Fish Introductions in Population Restoration	V-7
Stage II	<u>*</u>	
Stage III		
Founding new populations of threatened fishes		
	LITERATURE CITED	VI
GLOSSARY VII		
	GLOSSARY	VII

CHAPTER I

MOLECULAR GENETICS AND CYTOGENTICS

Generalized Cell

Cell membrane



Nucleus and Chromosomes

Introduction

Why be concerned about genetics?

From a genetics perspective organisms can be considered temporary carriers for a set of genes.

Genes not organisms are passed on from generation to generation.

Genetics is the study of how genes are inherited from one generation to the next and how they affect the characteristics of progeny.

What is a **gene**?

A specific segment of a molecule of **deoxyribonucleic acid (DNA)** to which a particular function can be assigned.

What is **DNA**?

A tightly coiled, helical molecule composed of:

phosphate

sugar: deoxyribose

four different nitrogenous bases

two purines: adenine (A) and guanine (G), 2 rings two pyrimidines: cytosine (C) and thymine (T), 1 ring carbon, nitrogen, hydrogen, oxygen

Nucleoside: base plus sugar

Nucleotide: nucleoside plus phosphate (Fig. 1)

DNA is composed of two strands of **complementary** nucleotides. That is, the sequence of one strand can always be converted to the sequence of the other because A always pairs with T and G with C (Fig.2).

<u>Complementarity</u> is very important. Allows for replication since given one strand you can form the other and allows DNA to act as an information storage and retrieval system.

Figure 2

•		
S 	 A====T	 S
P		P
S 	 G====C	 S
P		P
 S 	 C====G	 S
P		Р
 S 	 T====A	 S
Р 		Р
 S	 A====T	 S
		•

What is a **gene**?

A specific nucleotide sequence coding for the construction of a protein (**structural gene**) or that regulates the time and place of expression of other genes (**regulatory genes**).

Genes do not exist as individual units in the cell nucleus, but as a specific part of a chromosome.

What is a **chromosome**?

Single, tightly coiled molecule of DNA in association with protein constituting a linear array of genes.

Occur in pairs in practically all vertebrates.

Members of each pair usually carry the same set of genes because they are derived from a single common ancestor (**homologous**).

1 chromosome set: diploid (2N)
½ chromosome set: haploid (1N)
2 chromosome sets: tetraploid (4N)
½ chromosome sets: triploid (3N)

Usually divided into two classes:

sex chromosomes: have genes involved in sex determination **autosomes**: do not have genes involved in sex determination

Number of chromosomes (2N) is usually highly variable among species but is generally constant from generation to generation within a species (Table 1). 2N can be used for species detection and identification.

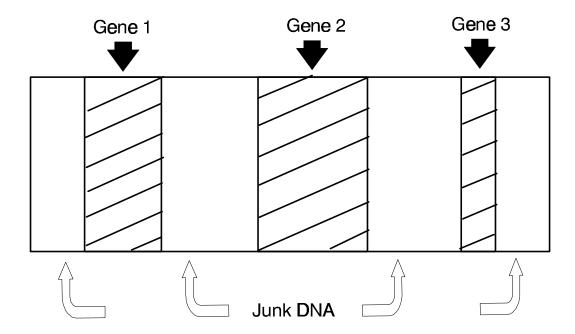
Table 1: Diploid number of chromosomes in madtom catfish (Gold et al. 1980).

Species	2N
Caddo	40
Frecklebelly	42
Tadpole	42
Least	46
Speckled	46
Freckled	48
Brindled	50
Margined	54

The linear array of genes on a chromosome is not continuous.

Adjacent genes are often interrupted by intervening sequences of DNA with no known function ("**Junk DNA**"; Fig. 3).

Figure 3



Genes themselves are often broken into pieces. There are four general parts of a gene (Fig. 4):

upstream flanking region: signals start of a gene

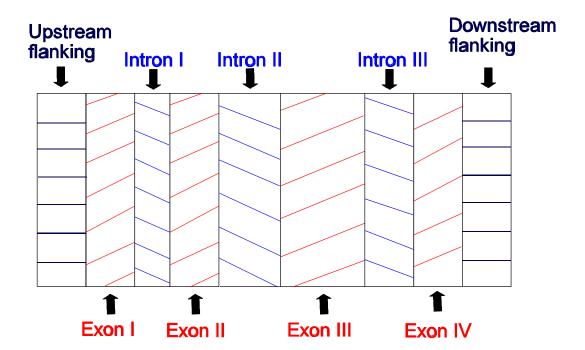
exons: code or perform gene function

introns: intervening sequences between exons

downstream flanking region: signals end of a gene

Introns with no known function are often interspersed among exons, hence genes in pieces.

Figure 4



Protein Construction

Proteins are composed of **amino acids** (Fig. 5). There are many kinds but only 20 are common.

<u>Polypeptide</u>: strand of amino acids formed by bonding of the carboxyl group of one to the amino group of the adjacent amino acid (**<u>peptide bond</u>**) (Fig. 6).

The type and position of each amino acid in a polypeptide represents the **primary structure** of protein.

Nucleotide sequence of a gene determines the primary structure of a protein.

Some polypeptides are proteins but not all proteins are a polypeptide.

Some proteins are composed of two or more polypeptides that may have the same or a different primary structure. Different polypeptides may be coded by the same gene or a different gene; 2 polypeptides from the \propto and 2 polypeptides from the \hat{a} hemoglobin genes combine to form hemoglobin.

Quaternary structure: number of polypeptides in a protein.

1 = monomer

2 = dimer

3 = trimer

4 = tetramer

How does gene structure determine protein primary structure?

There are three players in the game.

DNA = boss RNA = workers protein = product

What is **ribonucleic acid (RNA)**?

A molecule composed of:

phosphate sugar: ribose

four different nitrogenous bases

two purines: A and G

two pyrimidines: C and Uracil (U)

Different from DNA only because it is single stranded, has a different sugar, and U replaces T.

Generalized Amino Acid

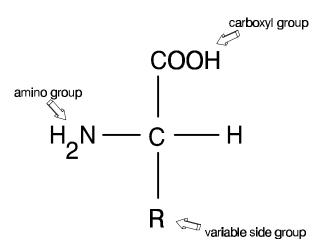
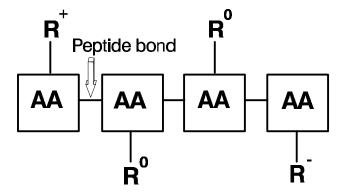


Figure 6



How is RNA made?

It is transcribed directly from a gene.

DNA "unzips" and one strand serves as a RNA template (Fig. 7). Strand transcribed varies from gene to gene but not within a gene.

A strand of RNA complementary to the DNA is made by an enzymatically controlled reaction.

DNA	RNA	
A	U	
G	C	
C	G	
A	U	

There are three basic kinds of RNA:

Ribosomal (rRNA): are part of the structure of ribosomes, cytoplasmic organelles attached to endoplasmic reticulum that serve as the site of protein synthesis (Cell Diagram, page I-2). Three kinds: 5S, 18S, 28S.

<u>Transfer (tRNA)</u>: bond to a specific amino acid and 'carry' to ribosomes for protein construction. Many kinds; at least one for every amino acid.

<u>Messenger (mRNA)</u>: brings complement of gene structure from the nucleus to the ribosome where its structure is translated by tRNA to construct a protein. Many, many kinds; at least one for every structural gene.

Figure 7

Growing mRNA strand ۲ TCG TTCAGCAACGGCT A C G G C T G C C G ∢ ⊢ O QQ \triangleleft \vdash GUUC/ CAA G O C B C Transcribed Untranscribed **DNA** strand

DNA strand

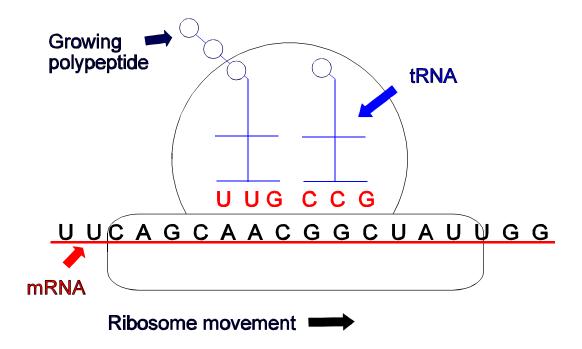
Back to protein construction.

mRNA leaves nucleus and attaches to ribosome in cytoplasm.

tRNA attaches to ribosome and aligns its three nucleotide <u>anticodon</u> region with the three nucleotide <u>codon</u> region of mRNA.

Amino acids of adjacent tRNA's are joined until entire mRNA message is translated into a polypeptide (Fig. 8).

Figure 8



Genetic code.

Sequence of three mRNA nucleotides (triplet or codon) that code for an amino acid.

Degenerate or redundant because most amino acids are coded by more than one triplet, e.g.,

mRNA codon	tRNA anticodon	amino acid
GUU GUA GUC	CAA CAU CAG	valine valine valine
UUA UUG	AAU AAC	leucine leucine
AUG	UAC	methionine
UGG	ACC	tryptophan

Only methionine and tryptophan have a single codon. Methionine codon initiates polypeptide synthesis.

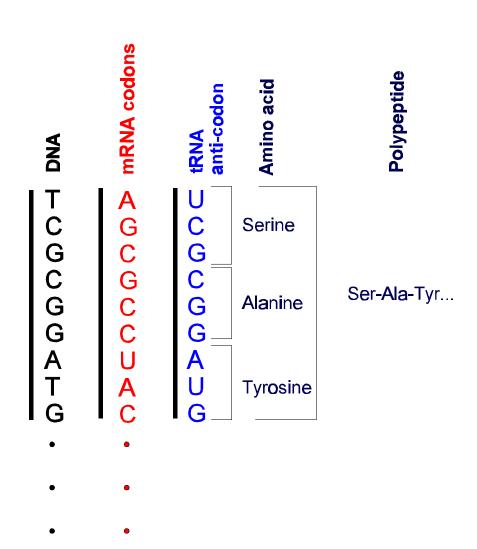
The code is nonoverlapping.

AGCGCCUAG

<u>AGC</u>	<u>GCC</u>	<u>UAG</u>
Codon 1	Codon 2	Codon 3
(serine)	(alanine)	(terminate synthesis)

Summary.

Chromosomes are large molecules mainly composed of a tightly coiled molecule of DNA in association with protein. Genes are a specific segment of DNA of a specific chromosome pair that code for the primary structure of proteins, formation of RNA, or regulate the time and place of expression of other genes. During protein synthesis DNA is transcribed to mRNA and translated to a polypeptide by tRNA on cytoplasmic ribosomes (Fig. 9).



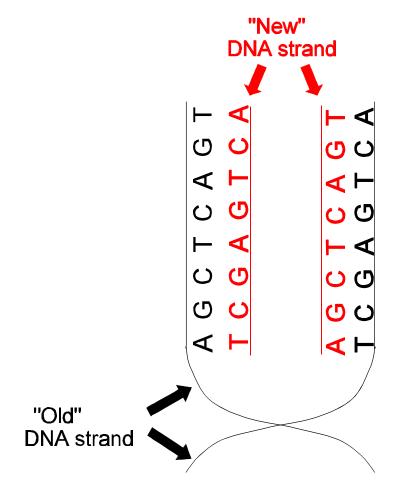
Cell Division

DNA replication.

Old strand of DNA serves as a template for construction of a new complementary strand by an enzymatically controlled reaction.

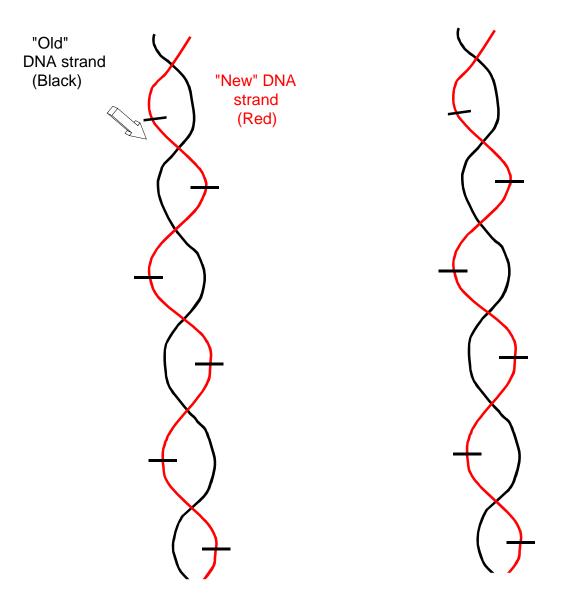
Double helix unzips at multiple sites and replication begins on both exposed strands (Fig. 10).

Figure 10



Semiconservative form of replication because each resulting molecule is half old and half new (Fig. 11).

Figure 11



Two basic types of cell division.

Mitosis: division for body growth, maintenance, and differentiation.

Meiosis: division for the production of **gametes**; eggs and sperm.

Mitosis

Problem: produce one cell from another so they are genetically identical.

Solution: one round of DNA replication followed by one cell division.

Interphase: period of protein synthesis and DNA replication (Fig. 12).

Once fully replicated the replicates or sister strands are physically attached at a region called the **centromere**.

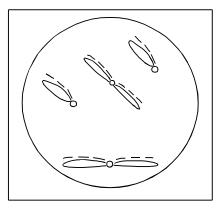
Prophase: chromosomes are fully replicated and begin moving to cell center or equator (Fig. 13).

<u>Metaphase</u>: chromosomes align along equator and spindle fibers attach to centromeres (Fig. 14).

Anaphase: sister chromosomes split apart at centromere and are carried to opposite poles by **spindle fibers** (Fig. 15).

<u>Telophase</u>: nuclear membrane reforms and cell divides in two (Fig. 16).

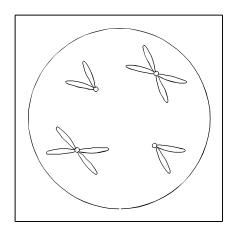
Interphase



Period of protein synthesis and DNA replication

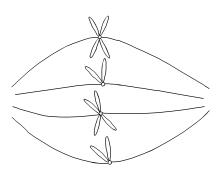
Figure 13

Prophase



Chromosomes are fully replicated and begin to move to cell center or equator

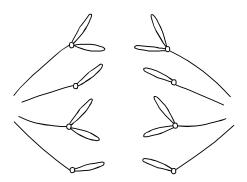
Metaphase



Chromosomes align along equator and spindle fibers attach to centromeres

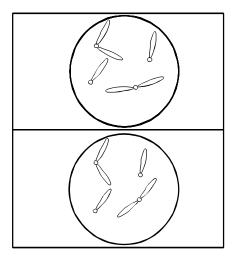
Figure 15

Anaphase



Sisters split at centromere and go to opposite poles

Telophase



Nuclear membrane reforms and cell divides in two

Meiosis

Problem: keep the amount of DNA constant from generation to generation or to produce a

haploid cell. That is, one having only 1N chromosomes.

Solution: one round of DNA replication followed by two cell divisions.

<u>Interphase</u>: protein synthesis and DNA replication.

Reduction Division

Prophase I: homologous pairs of sister chromosomes form and move to

equator. Note homologous pairs are not formed in mitosis

(Fig. 17).

Metaphase I: homologous pairs align along equator and spindle fibers attach to

centromeres (Fig. 18).

Anaphase I: members of each homologous pair move to opposite poles but

centromeres do not split (Fig. 19).

Telophase I: nuclear membrane reforms and members of homologous pairs are

in different cells but sisters are still attached at the centromere

(Fig. 20).

Equational Division

Prophase II: attached sisters in each cell begin to move to equator (Fig. 21).

Metaphase II: second alignment at equator and attachment of spindle fibers (Fig.

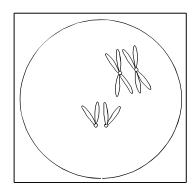
22).

Anaphase II: centromeres split and sisters move to opposite poles (Fig. 23).

Telophase II: nuclear membrane reforms and each resulting cell is haploid

(Fig. 24).

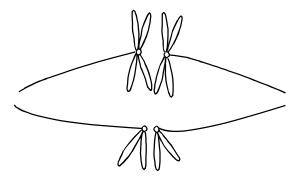
Prophase I



Chromosomes fully replicated, form <u>homologous pairs</u>, and move to the equator

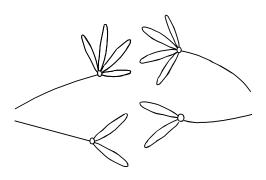
Figure 18

Metaphase I



Homologous pairs align along equator and spindle fibers attach to centromeres

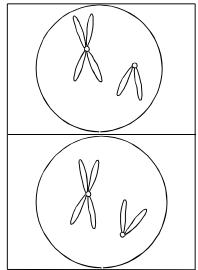
Anaphase I



Members of each homologous pair move to opposite poles: centromeres <u>do not</u> divide

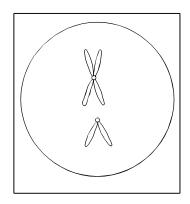
Figure 20

Telophase I



Nuclear membrane reforms but sisters still attached; homologous pairs broken up

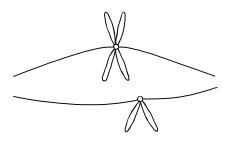
Prophase II



Second movement to equatorial plane

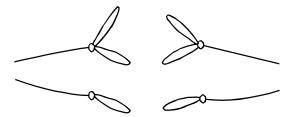
Figure 22

Metaphase II



Second alignment at equator and attachment of spindle fibers

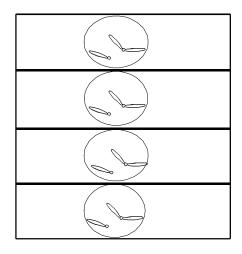
Anaphase II



Centromeres split and sisters move to opposite poles

Figure 24

Telophase II



Nuclear membrane reforms; each cell has only one copy of each pair of chromosomes (haploid)

Sperm Formation: cytoplasm is usually equally divided among daughter cells so each

precursor cell yields four sperm cells.

Egg Formation: cytoplasm is usually not equally divided among daughter cells so each

precursor yields one egg (almost all the original cytoplasm) and three **polar bodies** (have very little cytoplasm). It is energetically more costly

to be female.

Table 2: Contrast of Mitosis and Meiosis

Mitosis	Meiosis
1. Single DNA replication	1. Single DNA replication
2. One cell division	2. Two cell divisions
3. Homologous chromosomes do not pair	3. Homologous chromosomes pair
4. Centromeres divide	4. Centromeres do not divide until second division
5. Chromosome number maintained at 2N	5. Chromosome number halved to 1N
6. Produces two genetically identical cells	6. Produces four genetically different sperm cells or one egg and three polar bodies

Triploids and Gynogenetic Diploids

Green or unfertilized eggs have only completed the first meiotic division; i.e. they are diploid (2N).

Second meiotic division is completed only when egg is fertilized.

Second meiotic division can be prevented using heat shock or pressure.

<u>Triploid</u> (3N) is formed by using heat shock or pressure shortly after fertilization. Result is a diploid egg (2N) fertilized by a haploid sperm (1N). Is not 100% effective, some individuals are diploid. Thus, to release 100% triploids all fish must individually be checked for ploidy.

Gynogenetic diploid is a diploid individual whose chromosomes only came from the maternal parent (mother).

Formed by using sperm subjected to ultraviolet light (UV). UV fragments the DNA so functionally the sperm is 0N.

0N sperm is motile so is used to 'fertilize' eggs and heat shock or pressure shortly after fertilization results in a gynogenetic diploid; 2N egg plus 0N sperm.

Fish are highly inbred so do not constitute a good management tool but are a good research tool.